RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/576,982
Source:	IFWP.
Date Processed by STIC:	5/3/06

ENTERED



IFWP

DATE: 05/03/2006 RAW SEQUENCE LISTING PATENT APPLICATION: US/10/576,982 TIME: 11:21:20 Input Set : A:\50021-026.txt Output Set: N:\CRF4\05032006\J576982.raw 3 <110> APPLICANT: Queensland University of Technology 5 <120> TITLE OF INVENTION: Protein expression 7 <130> FILE REFERENCE: S80769691 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/576,982 C--> 9 <141> CURRENT FILING DATE: 2006-04-24 9 <150> PRIOR APPLICATION NUMBER: 2003905886 10 <151> PRIOR FILING DATE: 2003-10-24 12 <160> NUMBER OF SEQ ID NOS: 20 14 <170> SOFTWARE: PatentIn version 3.2 16 <210> SEQ ID NO: 1 17 <211> LENGTH: 50 18 <212> TYPE: PRT 19 <213> ORGANISM: Lactobacillus fermentum 21 <400> SEQUENCE: 1 23 Asp Thr Ile Tyr Thr Val Gln Ser Gly Asp Thr Leu Ser Gly Ile Ser 10 27 Tyr Lys Phe Ala Lys Asp Asn Ser Met Ile Asn Asp Leu Ala Lys Lys 20 25 31 Asn Asn Ile Gln Asp Ile Asn Lys Ile Phe Val Gly Gln Lys Leu Ile 40 35 Ile Lys 50 39 <210> SEQ ID NO: 2 40 <211> LENGTH: 81 41 <212> TYPE: PRT 42 <213> ORGANISM: Lactobacillus fermentum 44 <400> SEQUENCE: 2 46 Ser Tyr Thr Ser Asn Ala Ser Gly Ser Glu Ala Ala Ala Lys Ala Trp 50 Ile Ala Gly Arg Glu Ser Gly Gly Asn Tyr Asn Ala Thr Asn Gly Gln 20 54 Tyr Ile Gly Lys Tyr Gln Leu Ala Ala Ser Tyr Leu Gly Gly Asp Tyr 40 58 Ser Pro Ala Asn Gln Glu Arg Val Ala Asp Gln Tyr Val Ala Ser Arg 55 62 Tyr Gly Ser Trp Thr Ala Ala Gln Gln Phe Trp Gln Ala Asn Gly Trp 63 65 66 Tyr

73 <213> ORGANISM: Lactobacillus fermentum

70 <210> SEQ ID NO: 3 71 <211> LENGTH: 44 72 <212> TYPE: PRT

75 <400> SEQUENCE: 3

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77 Ser Asp Gly Glu Ile Gln Glu Tyr Asn Ala Gln Asn Ala Asn Ala 10 81 Asn Val Ala Asn Asn Asn Thr Gln Ala Thr Gln Gln Gln Thr Ala Gln 85 Ala Gln Pro Gln Gln Ala Gln Ser Gln Ala Asn Gln 35 89 <21.0> SEQ ID NO: 4 90 <211> LENGTH: 30 91 <212> TYPE: PRT 92 <213> ORGANISM: Lactobacillus fermentum 94 <400> SEQUENCE: 4 96 Met Ile Ser Lys Lys Asn Phe Ala Lys Val Ser Ala Thr Leu Gly Ala 10 5 100 Val Ala Leu Gly Val Ser Ala Thr Ala Thr Ala Ala Asn Ala 101 20 104 <210> SEQ ID NO: 5 105 <211> LENGTH: 175 106 <212> TYPE: PRT 107 <213> ORGANISM: Lactobacillus fermentum 109 <400> SEQUENCE: 5 111 Asp Thr Ile Tyr Thr Val Gln Ser Gly Asp Thr Leu Ser Gly Ile Ser 112 1 115 Tyr Lys Phe Ala Lys Asp Asn Ser Met Ile Asn Asp Leu Ala Lys Lys 20 25 119 Asn Asn Ile Gln Asp Ile Asn Lys Ile Phe Val Gly Gln Lys Leu Ile 40 123 Ile Lys Ser Asp Gly Glu Ile Gln Glu Tyr Asn Ala Gln Asn Ala Ala 55 127 Asn Ala Asn Val Ala Asn Asn Asn Thr Gln Ala Thr Gln Gln Gln Thr 70 131 Ala Gln Ala Gln Pro Gln Gln Ala Gln Ser Gln Ala Asn Gln Ser Tyr 85 90 135 Thr Ser Asn Ala Ser Gly Ser Glu Ala Ala Lys Ala Trp Ile Ala 105 139 Gly Arg Glu Ser Gly Gly Asn Tyr Asn Ala Thr Asn Gly Gln Tyr Ile 120 143 Gly Lys Tyr Gln Leu Ala Ala Ser Tyr Leu Gly Gly Asp Tyr Ser Pro 135 144 130 140 147 Ala Asn Gln Glu Arg Val Ala Asp Gln Tyr Val Ala Ser Arg Tyr Gly 150 155 151 Ser Trp Thr Ala Ala Gln Gln Phe Trp Gln Ala Asn Gly Trp Tyr 170 165 155 <210> SEQ ID NO: 6 156 <211> LENGTH: 205 157 <212> TYPE: PRT 158 <213> ORGANISM: Lactobacillus fermentum 160 <400> SEQUENCE: 6 162 Met Ile Ser Lys Lys Asn Phe Ala Lys Val Ser Ala Thr Leu Gly Ala 163 1 10

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166 167	Val	Ala	Leu	Gly 20	Val	Ser	Ala	Thr	Ala 25	Thr	Ala	Ala	Asn	Ala 30	Asp	Thr	
	Ile	Tyr	Thr 35		Gln	Ser	Gly	Asp		Leu	Ser	Gly	Ile 45		Tyr	Lys	
	Phe	Ala 50	Lys	Asp	Asn	Ser	Met 55		Asn	Asp	Leu	Ala 60		Lys	Asn	Asn	
			Asp	Ile	Asn	Lys.		Phe	Val	Gly	Gln 75		Leu	Ile	Ile	Lys 80	
182		Asp	Gly	Glu			Glu	Tyr	Asn			Asn	Ala	Ala			
183 186	Asn	Val	Ala	Asn	85 Asn	Asn	Thr	Gln	Ala	90 Thr	Gln	Gln	Gln	Thr	95 Ala	Gln	
187	77-	C1 m	Pro	100	C15	77-	C1 n	Cor	105	ח ד ה	7 an	Cl n	Cor	110	Thr	Cor	
191	AIA	GIII	115	GIII	GIII	AIa	GIII	120	GIII	AIa	ASII	GIII	125	TYL	1111	561	
	Asn		Ser	Gly	Ser	Glu		Ala	Ala	Lys	Ala		Ile	Ala	Gly	Arg	
195	~1	130	~1	~1··	7 an	Ma ess	135	717	Πb~	7 an	C1	140	TT	тло		Tara	
	145	ser	Gly	СТУ	ASII	151 150	ASII	Ala	THE	ASII	155	GIII	TAL	тте	GIY	LуS 160	
		Gln	Leu	Ala	Ala		Tvr	Tieu	Glv	Glv		Tvr	Ser	Pro	Ala		
203	-2,-			-	165		-		4	170	-	•			175		400
206	Gln	Glu	Arg	Val	Ala	Asp	Gln	Tyr	Val	Ala	Ser	Arg	Tyr	Gly	Ser	Trp	
207			-	180					185					190			
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211			195					200					205				
	214 <210> SEQ ID NO: 7																
	15 <211> LENGTH: 90 16 <212> TYPE: DNA																
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			EQUE														
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222	gtt	agtg	caa d	egge	tact	ge to	gctaa	atgci	t								90
225	<21	0 > S	EQ II	ои с	: 8												
			ENGT		50												
			YPE:		T = ==	1		e.								•	
			RGAN: EOUEI			LODA	CIII	18 16	erme	ıcum							
			~			a a	aataa	rtga	r ac	actti	tcad	gtai	tttc	tta .	caaai	tttgct	60
	_															aacaag	120
		-	ttg g		_	-	_	_		, ,				5			150
		_	EQII	_	_			•	-								
239	<21	1> L	ENGT	H: 1	32												
			YPE:														
			RGAN:			coba	cill	ıs f	erme	ntum							
			EQUEI														
																gcaaac	60 120
			acc a		Laca	a do	Jaaci	aadCl	L gc	Luad	ycac	aac	CLCA	aca	ayca	caaagc	132
			EQ II		: 10												1,2
			ENGT														

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PATENT APPLICATION: US/10/576,982 TIME: 11:21:20

Input Set : A:\50021-026.txt

Output Set: N:\CRF4\05032006\J576982.raw

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   259 gaatcaggtg gtaactacaa cgccacaaac ggtcaataca ttggtaagta ccaattagct
                                                                              120
   261 gcatcatacc ttggtggtga ctactcacca gctaaccaag aacgcgttgc tgaccaatac
                                                                              180
263 gttgczagte gttacggtte ttggaetget geccaacaat tetggeaage aaacggttgg
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   265 tactaa
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   271 <213> ORGANISM: Lactobacillus fermentum
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   276 aaagacaaca gtatgatcaa tgatcttgct aagaagaaca atattcaaga tattaacaag
                                                                              120
   278 atttttgttg gtcaaaagtt aatcatcaag agcgatggtg aaattcaaga atacaatgct
                                                                              180
   280 caaaatgcag ctaatgcaaa tgtagcaaac aacaatactc aagctacaca acaacaaact
                                                                              240
   282 getcaageac aacetcaaca ageacaaage caagetaace aaagetacae tteaaatget
	imes 284 	imes 	ext{caggtteag} (aagetgetge taaagettgg attgeeggte gtgaat	ilde{e}aage tggtaactac
                                                                              360-
   286 aacgccacaa acggtcaata cattggtaag taccaattag ctgcatcata ccttggtggt
                                                                              420
   288 gactactcac cagctaacca agaacgcgtt gctgaccaat acgttgcaag tcgttacggt
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   298 <400> SEQUENCE: 12
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   301 gttagtgcaa cggctactgc tgctaatgct gacactatct acaccgtaca aagtggtgac
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   303 acactttcag gtatttctta caaatttgct aaagacaaca gtatgatcaa tgatcttgct
                                                                              180
   305 aagaagaaca atattcaaga tattaacaag atttttgttg gtcaaaagtt aatcatcaag
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   307 agcgatggtg aaattcaaga atacaatgct caaaatgcag ctaatgcaaa tgtagcaaac
                                                                              300
   309 aacaatactc aagctacaca acaacaaact qctcaaqcac aacctcaaca agcacaaagc
                                                                              360
   311 caagetaace aaagetacae tteaaatget teaggtteag aagetgetge taaagettgg
                                                                              420
   313 attgccggtc gtgaatcagg tggtaactac aacgccacaa acggtcaata cattggtaag
                                                                              480
                                                                              540
   315 taccaattag ctgcatcata ccttggtggt gactactcac cagctaacca agaacgcgtt
   317 getgaccaat aegttgeaag tegttaeggt tettggaetg etgeecaaca attetggeaa
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334 <221> NAME/KEY: misc_feature 335 <222> LOCATION: (24)..(24)

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395 <222> LOCATION: (63)..(63)

396 <223> OTHER INFORMATION: any nucleotide

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 05/03/2006
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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VERIFICATION SUMMARYDATE: 05/03/2006PATENT APPLICATION: US/10/576,982TIME: 11:21:21

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